

Entrez PubMed Nucleotide Protein Genome Structure PMC Taxonomy

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Entrez Nucleotide

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☐ 1: [Reports](#) [Links](#)
[NM_078456](#)

Caenorhabditis elegans ceramide glucosyltransferase / glucosylceramide synthase family member (50.1 kD) (XS315) mRNA
gi|25147525|ref|NM_078456.2|[[25147525]]

Related resources

☐ 2: [Reports](#) [Links](#)
[NM_127546](#)

Arabidopsis thaliana ceramide glucosyltransferase, putative (At2g19880) mRNA, complete cds
gi|18399183|ref|NM_127546.1|[[18399183]]

☐ 3: [Reports](#) [Links](#)
[XM_538788](#)

PREDICTED: Canis familiaris similar to Ceramide glucosyltransferase (Glucosylceramide synthase) (GCS) (UDP-glucose:N-acylsphingosine D-glucosyltransferase) (UDP-glucose ceramide glucosyltransferase) (GLCT-1) (LOC481666), mRNA
gi|57094232|ref|XM_538788.1|[[57094232]]

☐ 4: [Reports](#) [Links](#)
[NW_139871](#)

Canis familiaris chromosome 11 genomic contig, whole genome shotgun sequence
gi|54126007|ref|NW_139871.1|Cfa11_WGA36_1[[54126007]]

☐ 5: [Reports](#) [Links](#)
[NM_031795](#)

Rattus norvegicus UDP-glucose ceramide glucosyltransferase (Ugcg), mRNA
gi|25742622|ref|NM_031795.1|[[25742622]]

☐ 6: [Reports](#) [Links](#)
[NM_004775](#)

Homo sapiens UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase,

polypeptide 6 (B4GALT6), mRNA
gi|13929471|ref|NM_004775.2|[13929471]

☐ **7:** Reports Links
NM_003358

Homo sapiens UDP-glucose ceramide glucosyltransferase (UGCG), mRNA
gi|4507810|ref|NM_003358.1|[4507810]

☐ **8:** Reports Links
NT_086888

Homo sapiens chromosome 18 genomic contig, alternate assembly
gi|51474639|ref|NT_086888.1|Hs18_86553[51474639]

☐ **9:** Reports Links
NT_086754

Homo sapiens chromosome 9 genomic contig, alternate assembly
gi|51467694|ref|NT_086754.1|Hs9_86435[51467694]

☐ **10:** Reports Links
NT_008470

Homo sapiens chromosome 9 genomic contig
gi|51467683|ref|NT_008470.17|Hs9_8627[51467683]

☐ **11:** Reports Links
NT_010966

Homo sapiens chromosome 18 genomic contig
gi|29823167|ref|NT_010966.13|Hs18_11123[29823167]

☐ **12:** Reports Links
NC_003071

Arabidopsis thaliana chromosome 2, complete sequence
gi|30698031|ref|NC_003071.3|[30698031]

☐ **13:** Reports Links
NC_003284

Caenorhabditis elegans chromosome X, complete sequence
gi|32566600|ref|NC_003284.3|[32566600]

☐ **14:** Reports Links
AJ224156

Rattus norvegicus mRNA for ceramide glucosyltransferase
gi|2924339|emb|AJ224156.1|RNAJ4156[2924339]

☐ **15:** Reports Links
AF367245

Gossypium arboreum ceramide glucosyltransferase mRNA, complete cds
gi|14718994|gb|AF367245.1|AF367245[14718994]


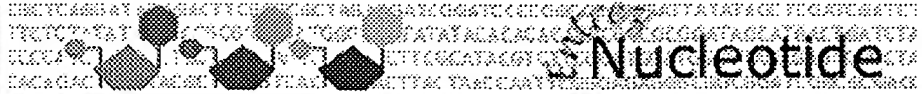

- ☐ **16:**
AF364403 Reports Links
Pichia pastoris ceramide glucosyltransferase gene, complete cds
gi|14718992|gb|AF364403.1|AF364403[14718992]
- ☐ **17:**
AF364402 Reports Links
Magnaporthe grisea ceramide glucosyltransferase mRNA, complete cds
gi|14718990|gb|AF364402.1|AF364402[14718990]
- ☐ **18:**
AF364401 Reports Links
Caenorhabditis elegans ceramide glucosyltransferase mRNA, complete cds
gi|14718988|gb|AF364401.1|AF364401[14718988]
- ☐ **19:**
AF119416 Reports Links
Mus musculus GM3 synthase mRNA, complete cds
gi|7670072|gb|AF119416.1|AF119416[7670072]
- ☐ **20:**
AF119415 Reports Links
Homo sapiens GM3 synthase mRNA, complete cds
gi|7670070|gb|AF119415.1|AF119415[7670070]

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Reports

[Links](#)

Homo sapiens UDP-Gal:glucosylceramide beta-1,4-galactosyltransferase mRNA, complete cds
gi|5305556|gb|AF097159.1|AF097159[5305556]

☐ 22:[AF047707](#)

Reports

[Links](#)

Rattus norvegicus UDP-glucose:ceramide glycosyltransferase mRNA, complete cds
gi|4105566|gb|AF047707.1|AF047707[4105566]

☐ 23:[AF097158](#)

Reports

[Links](#)

Mus musculus UDP-Gal:glucosylceramide beta-1,4-galactosyltransferase mRNA, complete cds
gi|5305554|gb|AF097158.1|AF097158[5305554]

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